

SEQUENCE LISTING

<110> COHEN, STEPHEN
BOUWMEESTER, ANTONIUS
ROYET, JULIEN

<120> REGULATOR OF NOTCH SIGNALING ACTIVITY

<130> 55880(71745)

<140> 09/830,980

<141> 2001-05-02

<150> PCT/IB99/01891

<151> 1999-11-03

<150> GB 9824045.0

<151> 1998-11-03

<160> 16

<170> PatentIn Ver. 2.1

<210> 1

<211> 480

<212> PRT

<213> Drosophila sp.

<400> 1

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Pro	Ala	Gly	Ile	Thr	Thr	Gln	Gln	Leu	Gly	Leu	Ile	Cys	Asn	Ala	Leu
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Glu	Ile	Lys	Lys	Ser	Leu	Glu	Asp	Thr	Leu	Asp	Leu	Ala	Ser	Val	Asp
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Gly	Ser	Gly	Asp	Thr	Thr	Val	Arg	Leu	Trp	Asp	Leu	Asn	Thr	Glu	Thr
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Pro His Phe Thr Cys Thr Gly His Lys Gln Trp Val Leu Cys Val Ser
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 Ile Ile Ile Trp Asp Pro Glu Thr Gly Gln Gln Lys Gly Arg Pro Leu
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 Ser Gly His Lys Lys His Ile Asn Cys Leu Ala Trp Glu Pro Tyr His
 195 200 205
 Arg Asp Pro Glu Cys Arg Lys Leu Ala Ser Ala Ser Gly Asp Gly Asp
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 Cys Arg Ile Trp Asp Val Lys Leu Gly Gln Cys Leu Met Asn Ile Ala
 225 230 235 240
 Gly His Thr Asn Ala Val Thr Ala Val Arg Trp Gly Gly Ala Gly Leu
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 Ile Tyr Thr Ser Ser Lys Asp Arg Thr Val Lys Met Trp Arg Ala Ala
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 Asp Gly Ile Leu Cys Arg Thr Phe Ser Gly His Ala His Trp Val Asn
 275 280 285
 Asn Ile Ala Leu Ser Thr Asp Tyr Val Leu Arg Thr Gly Pro Phe His
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 Ala Ser Phe Asp Lys Ser Val Arg Leu Trp Arg Ala Ser Asp Gly Gln
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 Tyr Met Ala Thr Phe Arg Gly His Val Gln Ala Val Tyr Thr Val Ala
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<212> DNA

<213> *Drosophila* sp.

<400> 2

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<211> 513

<212> PRT

<213> *Saccharomyces cerevisiae*

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 35 40 45

Leu Arg Val Pro Gly Ala Ile Ser Glu Lys Gln Leu Glu Glu Leu Leu
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 Asn Gln Leu Asn Gly Thr Ser Asp Asp Pro Val Pro Tyr Thr Phe Ser
 65 70 75 80
 Cys Thr Ile Gln Gly Lys Lys Ala Ser Asp Pro Val Lys Thr Ile Asp
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 Ile Thr Asp Asn Leu Tyr Ser Ser Leu Ile Lys Pro Gly Tyr Asn Ser
 100 105 110
 Thr Glu Asp Gln Ile Thr Leu Leu Tyr Thr Pro Arg Ala Val Phe Lys
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 Thr Ile Leu Cys Ser Ala Phe Ala Pro His Thr Ser Ser Arg Met Val
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 165 170 175
 Thr Pro Met His Thr Leu Lys Gly His Tyr Asn Trp Val Leu Cys Val
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 Thr Ile Arg Leu Trp Asp Pro Lys Ser Gly Gln Cys Leu Gly Asp Ala
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 260 265 270
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 Thr Thr Leu Lys Val Trp Asp Val Arg Thr Arg Lys Leu Ser Val Asp
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<210> 4
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 <212> PRT
 <213> Codonanthus elegans

<220>
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 35 40 45
 Ser Gly Ala Glu Ile Val Asp Ser Ile Arg Lys Ser Leu Glu Glu Ile
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Asp Phe Glu Thr Leu Lys Leu Val Tyr Gln Pro Gln Ala Val Phe Arg
 65 70 75 80
 Val Arg Pro Val Thr Arg Cys Ser Ala Ser Ile Pro Gly His Gly Glu
 85 90 95
 Pro Val Ile Ser Ala Gln Phe Ser Pro Asp Gly Arg Gly Leu Ala Ser
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 Gly Ser Gly Asp Gln Thr Met Arg Ile Trp Asp Ile Glu Leu Glu Leu
 115 120 125
 Pro Leu His Thr Cys Lys Ser His Lys Ser Trp Val Leu Cys Ile Ala
 130 135 140
 Trp Ser Pro Asp Ala Thr Lys Ile Ala Ser Ala Cys Lys Asn Gly Glu
 145 150 155 160
 Ile Cys Ile Trp Asn Ala Lys Thr Gly Glu Gln Ile Gly Lys Thr Leu
 165 170 175
 Lys Arg His Lys Gln Trp Ile Xaa Xaa Leu Ala Trp Gln Pro Thr Val
 180 185 190
 Lys Met Trp Arg Ala Asp Asp Gly Val Met Cys Arg Asn Met Thr Gly
 195 200 205
 His Ala His Trp Ile Asn Thr Leu Ala Leu Asn Thr Asp Tyr Ala Leu
 210 215 220
 Arg Thr Ser Cys Phe Glu Pro Ser Lys Ile Asn Arg Met Thr Gly His
 225 230 235 240
 Met Gln Leu Val Asn Gln Val Val Phe Ser Pro Asp Thr Arg Tyr Ala
 245 250 255
 Ser Ala Ser Phe Asp Lys Ser Val Lys Leu Trp Cys Gly Arg Thr Gly
 260 265 270
 Lys Tyr Leu Ala Ser Phe Arg Gly His Val Gly Pro Val Tyr Gln Val
 275 280 285
 Ala Trp Ser Ala Asp Ser Arg Leu Leu Val Ser Gly Ser Ala Asp Ser
 290 295 300
 Thr Leu Lys Val Phe Glu Leu Lys Thr Lys Ser Leu Tyr Tyr Asp Leu
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 Pro Gly His Gly Asp Glu Val Phe Thr Val Asp Trp Ser Pro Glu Gly
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 Thr Lys Val Val Ser Gly Gly Lys Asp Lys Val Leu Lys Leu Trp
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<211> 103

<212> PRT
 <213> Mus sp.

<220>
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 <222> (39)
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 35 40 45
 Glu Glu Pro Leu Pro Leu Ala Phe Tyr Val His Asp Ala Glu Ile Val
 50 55 60
 Ser Ser Leu Gly Lys Thr Leu Glu Ser Gln Ser Val Glu Thr Glu Lys
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 Ile Val Asp Ile Ile Tyr Gln Pro Gln Ala Val Phe Arg Val Arg Ala
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 Val Thr Arg Cys Thr Ser Ser
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<210> 6
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 <212> PRT
 <213> Homo sapiens

<220>
 <221> MOD_RES
 <222> (66)
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 20 25 30
 Phe Phe Val His Asp Ala Glu Ile Val Ser Ser Leu Gly Lys Thr Leu
 35 40 45
 Glu Ser Gln Ala Val Glu Thr Glu Lys Val Leu Asp Ile Tyr Gln Pro
 50 55 60
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<210> 7
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 <212> PRT
 <213> *Xenopus laevis*

<400> 7

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Asp	Lys	Leu	Gln	Leu	Val	Cys	Asn	Ala	Leu	Leu	Gln	Glu	Glu	Asp	Pro	35	40	45	
Val	Pro	Leu	Ala	Phe	Phe	Val	Gln	Asp	Leu	Glu	Ile	Val	Thr	Ser	Leu	50	55	60	
Asp	Lys	Thr	Leu	Glu	Lys	Gln	Ser	Val	Glu	Thr	Glu	Lys	Val	Ile	Asp	65	70	75	80
Ile	Ile	Tyr	Gln	Pro	Gln	Ala	Val	Phe	Lys	Val	Arg	Ala	Val	Thr	Arg	85	90	95	
Cys	Thr	Ser	Ser	Leu	Glu	Gly	His	Thr	Glu	Ala	Val	Ile	Ser	Val	Ala	100	105	110	
Phe	Ser	Pro	Thr	Gly	Lys	Tyr	Leu	Ala	Ser	Gly	Ser	Gly	Asp	Thr	Thr	115	120	125	
Val	Arg	Phe	Trp	Asp	Leu	Ser	Thr	Glu	Thr	Pro	His	Phe	Thr	Ser	Lys	130	135	140	
Gly	His	Thr	His	Trp	Val	Leu	Ser	Ile	Ala	Trp	Ser	Pro	Asp	Gly	Lys	145	150	155	160
Lys	Leu	Ala	Ser	Gly	Cys	Lys	Asn	Ser	Gln	Ile	Phe	Ile	Trp	Asp	Pro	165	170	175	
Ser	Thr	Gly	Lys	Gln	Ile	Gly	Lys	Pro	Leu	Thr	Gly	His	Ser	Lys	Trp	180	185	190	
Ile	Thr	Trp	Leu	Cys	Trp	Glu	Pro	Leu	His	Leu	Asn	Pro	Glu	Ser	Arg	195	200	205	
Tyr	Leu	Ala	Ser	Ala	Ser	Lys	Asp	Cys	Thr	Ile	Arg	Ile	Trp	Asp	Thr	210	215	220	
Val	Met	Gly	Gln	Cys	Gln	Lys	Ile	Leu	Thr	Ser	His	Thr	Gln	Ser	Val	225	230	235	240
Thr	Ala	Val	Lys	Trp	Gly	Gly	Asp	Gly	Leu	Leu	Tyr	Ser	Ser	Ser	Gln	245	250	255	
Asp	Arg	Thr	Ile	Lys	Ala	Trp	Arg	Ala	Gln	Asp	Gly	Val	Leu	Cys	Arg	260	265	270	

Thr Leu Gln Gly His Ala His Trp Val Asn Thr Met Ala Leu Ser Thr
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 Asp Tyr Val Leu Arg Lys Gly Ala Phe Asn Pro Ala Asp Ala Ser Val
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 Asn Pro Gln Asp Met Ser Gly Ser Leu Glu Val Leu Lys Glu Lys Ala
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 Leu Lys Arg Ser Asn Glu Val Arg Gly Gln Gly Pro Glu Arg Leu Val
 325 330 335
 Ser Gly Ser Glu Asp Phe Thr Leu Phe Leu Trp Ala Pro Ala Glu Glu
 340 345 350
 Lys Lys Pro Leu Gln Arg Met Thr Gly His Gln Ala Leu Ile Asn Glu
 355 360 365
 Val Leu Phe Ser Pro Asp Thr Arg Ile Ile Ala Ser Ala Ser Phe Asp
 370 375 380
 Lys Ser Ile Lys Leu Trp Asp Gly Lys Thr Gly Lys Phe Leu Thr Ser
 385 390 395 400
 Leu Arg Gly His Val Ser Ala Val Tyr Gln Ile Ala Trp Ser Ala Asp
 405 410 415
 Ser Arg Leu Leu Val Ser Gly Ser Ser Asp Ser Thr Leu Lys Val Trp
 420 425 430
 Asp Ser Lys Thr Lys Lys Leu Leu Ile Asp Leu Pro Gly His Ala Asp
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 Glu Val Tyr Ser Val Asp Trp Ser Pro Asp Gly Gln Arg Val Ala Ser
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<210> 8

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: HA epitope

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<210> 9

<211> 149

<212> DNA
 <213> Artificial Sequence

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 <223> Description of Artificial Sequence: Primer

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 gccacgccac atacgatata ggcgcgcca 149

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<220>
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<210> 11
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 <223> Description of Artificial Sequence: Primer

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 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Conserved
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<210> 14
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 <213> Artificial Sequence

<220>
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 <212> DNA
 <213> Artificial Sequence

<220>
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<210> 16
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<220>
<223> Description of Artificial Sequence: Primer

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21